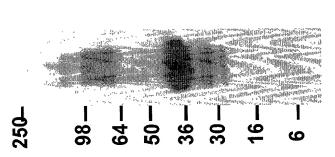
FIG. 1.

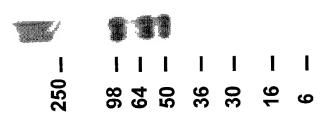
Multiple Alignment:

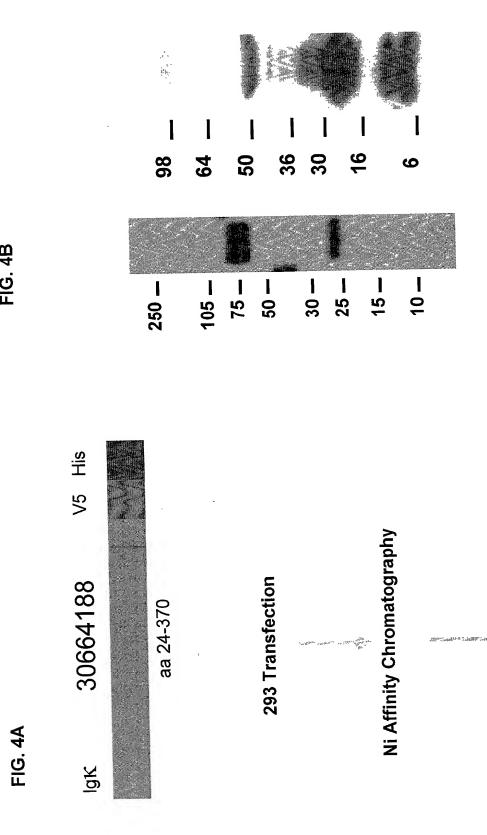
8%	119 114	179 165	238 214	296 274	356 331	33
FVYTLÍCANFCSCRDT SATPOSASIKALRNANLRRDESNHÍTDLYR RÍÐETIQWKG FGLLLÍFTSALAGOROGÍQAESNLSSKFOFSSNKBONGVODPO.HERIITWST	S PRFP N SYPRN L DLTWRLHS - QENTR IQLYFDNQFGLEEAENDICR YDFVEV EGI S PRFP HTYPRNT WLYWRLYAWENVWIQLTFDERFGLEDPEDDICK YDFVEV EEP	I IRBRWCGHKEVPPRIKIKSKINQIKIT FKSDDVFVAKPGFKIYVSIL EDFOPAAAS TI LGRWCGSGTVPGKQISKGNOIRIRFVSDEVFPSEPGFCIHVNAVMP	SVTSSTSGVSYNSPSVTDP-TLIADALDKKIAEFDTVEDLIKVFNP ESWOEDLEN QFTEAWSPSVLPPSALPLDLINNAITAFSTEDLENTEP ERWOLDLED	PRYRGESYHD. RKS. KWDIDRINDEAKRYSCTPRNYSYNIREELKL ANV WERPRC WOLLGKAFYFGRKSRVWDINLITEEVRLYSCTPRNESYSIREELKR TDT HEWPGD	CGGNC GOCTVNWRSOTONSGKTVKKYHEVLOFEPGHIKRRGRÅKTMALVDFOLDH CGGNC KCLHNCNEDODVPSKVTKKYHEVLOLRPKTGVRÆLHK SLTDKALEH	OCTOSSRPR (SEQ ID NO.2)
MIRLI 1 MS	NGY WO NGS PH	SHT ST Section 1	TNWE	MWLDTI LVRPI	LLW ORO	HERODO HEECDO
			[五]			
→ →	61 55	120 115	180 166	239 215	23	337
30664188.0.99 VEGF-E	30664188.0.99 VEGF-E	30664188.0.99 VEGF-E	30664188.0.99 180 ETINWE VEGF-E 166	30664188.0.99 VEGF-E	30664188.0.99 297 VEGF-E 275	30664188.0.99 357 VEGF-E 332

DESCRIPT

FIG. 3.

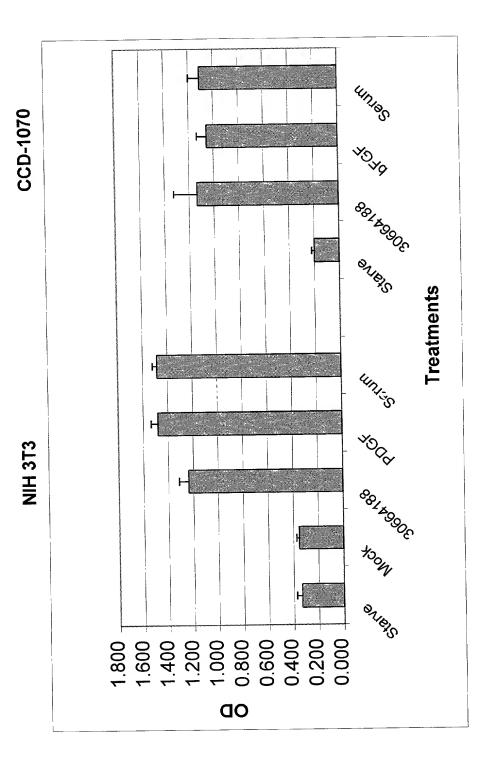




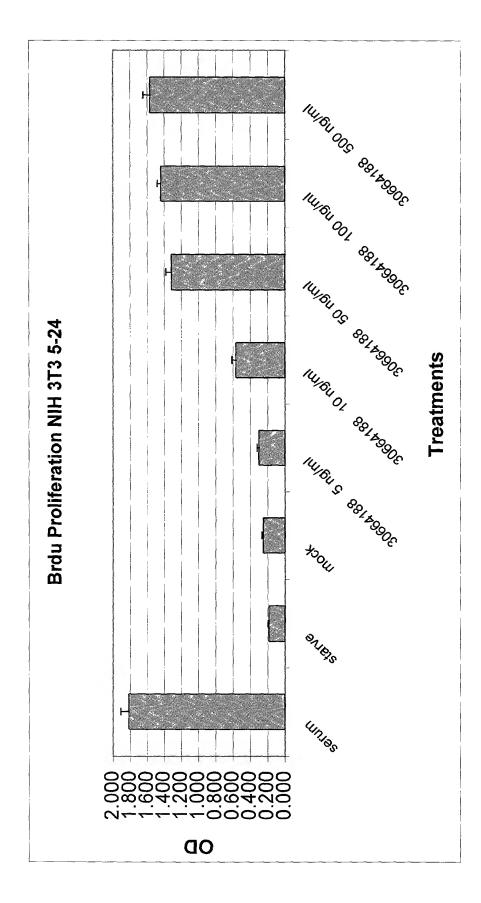


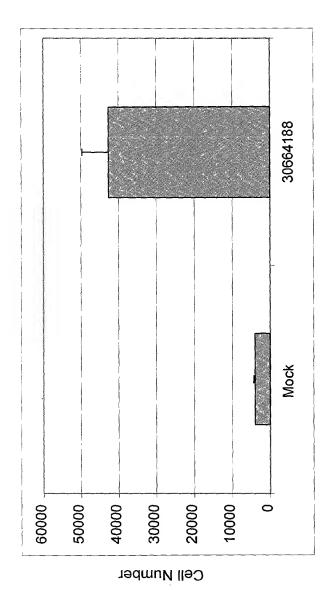
Imidazole Elution



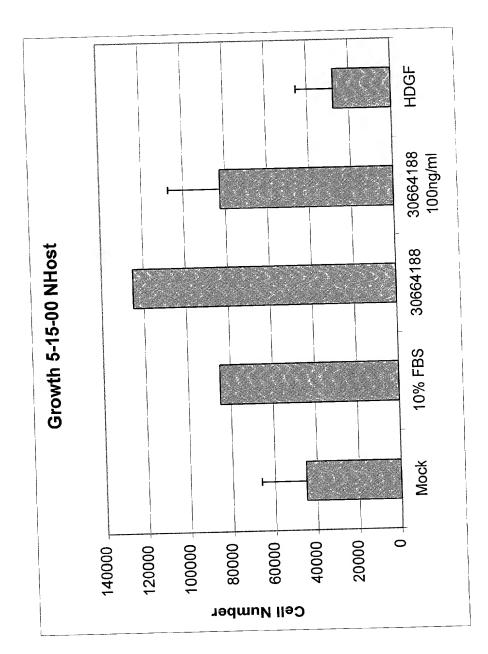




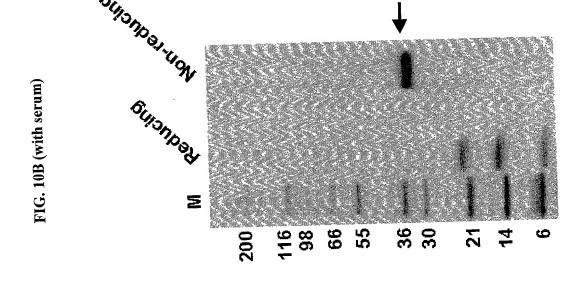




pCEP4sec(30664188) CM pCEP4sec CM



ppppe 1417



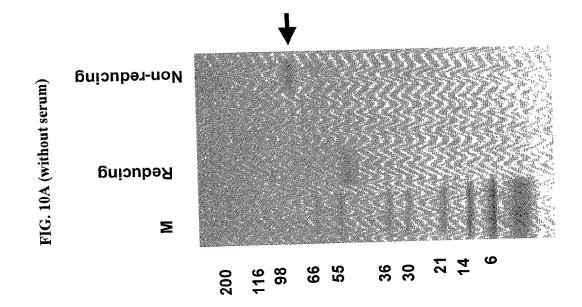
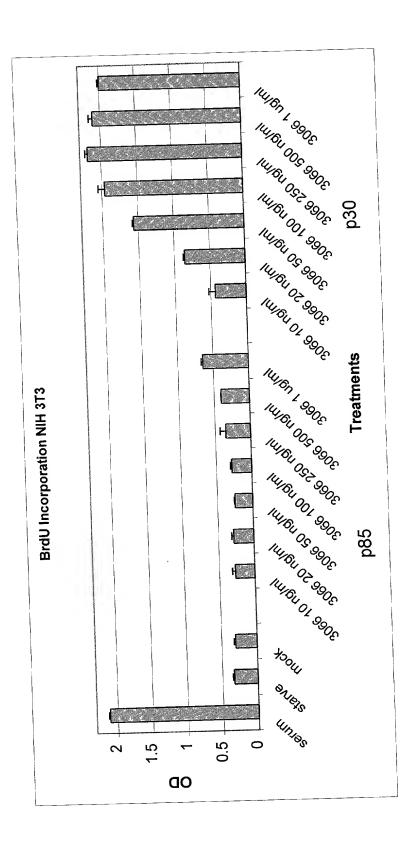


FIG. 10.

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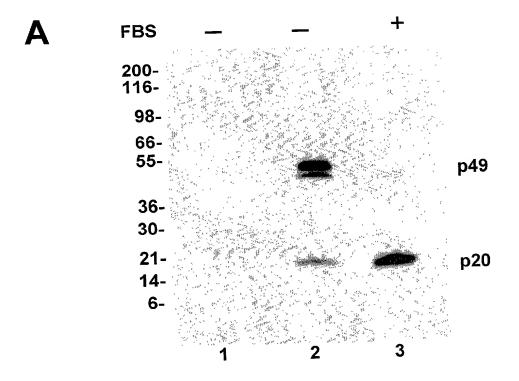


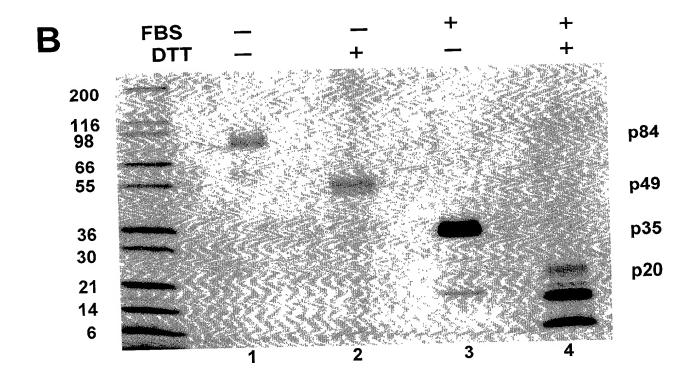
CTPRNYSVNI-REELKLANVVF--FPROLLVQROGGONCGCGTVNWRSCTC hPDGF D mPDGF D CTPRNHSVNL-REELKLTNAVF--FPRCLLVQRCGGNCGCGTVNWKSCTC PDGF C CTPRNFSVSI-REELKRTDTIF--WPGCLLVKRCGGNCACCLHNCNECQC CKTRTEVFEISRRLIDRTNANFLVWPPCVEVQRCSG---CCNNRNVQCRP PDGF B PDGF A CKTRTVIYEIPRSQVDPTSANFLIWPPCVEVKRCTG---CCNTSSVKCQP hPDGF D NS---GKTVKKYHEVLQFEPGHIKRRGRAKTMALVDIQLDHHER (SEQ ID NO:15) mPDGF D SS---GKTVKKYHEVLKFEPGHFKRRGKAKNMALVDIQLDHHERCDC (SEQ ID NO:16) VP---SKVTKKYHEVLQLRPKTGVRGLH-KSLTDVA--LEHHEECDC (SEQ ID NO:17) PDGF C TQVQLRPVQVRKIEIVRKKPIF----KKAT-VT----LEDHLACKC (SEQ ID NO:18) PDGF B SRVHHRSVKVAKVEYVRKKPKL----KEVQ-VR----LEEHLECAC (SEQ ID NO:19) PDGF A

 $\textbf{1} \overset{Exon \, 1}{\texttt{CGCAGGGCGGCGGCGGCGGGAGCAGAACCCGGCTTTTTCTTGGAGCGACGCTGTCTCTAGTCGCTGATCCCA}}$ aatglaccggctcatctttgtctacactctaatctgcgcaaacttttgcagctgtcgggacacttctgcaaccccgcaga MHRLIFVYTLICANFCSCRDTSATPQS Exonl Exon2 GCGCATCCATCAAAGCTTTGCGCAACGCCAACCTCAGGCGAGATGAGAGCAATCACCTCACAGACTTGTACCGAAGAGAT A S I K A L R N A N L R R D E S N H L T D L Y R R D GAGACCATCCAGGTGAAAGGAAACGGCTACGTGCAGAGTCCTAGATTCCCGAACAGCTACCCCAGGAACCTGCTCCTGAC 241 E T I Q V K G N G Y V Q S P R F P N S Y P R N L L L T W R L H S Q E N T R I Q L V F D N Q F G L E E A E N D Exon 2 Exon 3 ATATCTGTAGTATGATTTTGTGGAAGTTGAAGATATATCCGAAACCAGTACCATTATTAGAGGACGATGGTGTGGACAC ICRYDFVEVEDISETSTIIRGRWCGH AAGGAAGTTCCTCCAAGGATAAAATCAAGAACGAACCAAATTAAAATCACATTCAAGTCCGATGACTACTTTGTGGCTAA KEVPPRIKSRTNQIKITFKSDDYFVAK ACCTGGATTCAAGATTTATTATTCTTTGCTGAAGATTTCCAACCCGCAGCAGCTTCAGAGACCAACTGGGAATCTGTCA P G F K I Y Y S L L E D F Q P A A A S E T N W E S V T Exon 4 Exon 5 CAAGCTCTATTTCAGGGGTATCCTATAACTCTCCATCAGTAACGGATCCCACTCTGATTGCGGATGCTCTGGACAAAAAA S S I S G V S Y N S P S V T D P T L I A D A L D K K ${\tt ATTGCAGAATTTGATACAGTGGAAGATCTGCTCAAGTACTTCAATCCAGAGTCATGGCAAGAAGATCTTGAGAATATGTA}$ IAEFDTVEDLLKYFNPESWQEDLENMY
Exon 5 TCTGGACACCCCTCGGTATCGAGGCAGGTCATACCATGACCGGAAGTCAAAAGTTGACCTGGATAGGCTCAATGATGATG LDTPRYRGRSYHDRKSKVDLDRLNDDA 881 CCAAGCGTTACAGTTGCACTCCCAGGAATTACTCGGTCAATATAAGAGAAGAGCTGAAGTTGGCCAATGTGGTCTTCTTT K R Y S C T P R N Y S V N I R E E L K L A N V V F F CCACGTTGCCTCCTCGTGCAGCGCTGTGGAGGAAATTGTGGCTGTGGAACTGTCAACTGGAGGTCCTGCACATGCAATTC PRCLLVQRCGGNCGCGTVNWRSCTCNS AGGGARAACCGTGAAAAAGTATCATGAGGTATTACAGTTTGAGCCTGGCCACATCAAGAGGAGGGGTAGAGCTAAGACCA G K T V K K Y H E V L Q F E P G H I K R R G R A K T M TGGCTCTAGTTGACATCCAGTTGGATCACCATGAACGATGTGATTGTATCTGCAGCTCAAGACCACCTCGATAASAGAAT A L V D I Q L D H H E R C D C I C S S R P P R (SEO ID NO:20) 1201 GTGCACATCCTTACATTAAGCCTGAAAGAACCTTTAGTTTAAGGAGGGTGAGATAAGAGACCCTTTTCCTACCAGCAACC 1281 AAACTTACTACCAGCCTGCAATGCAATGAACACAAGTGGTTGCTGAGTCTCAGCCTTGCTTTAATGCCATGGCAAGT 1521 CTTACATATGTTTGAGTTAGACTCTTAAAATCCTTTGCCAAAATAAGGGATGGTCAAATATATGAAACATGTCTTTAGAA 1681 ATATTAAAAATCAAAAGATGAGGCTTTCTTACATATACATCTTAGTTG (SEO ID NO:21)

remain i

FIG. 14





A. 27 KDa

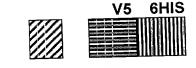
GRSYHDRKSKVDLDRL

GRSYHDRKSKVD..... (SEQ ID NO:23)

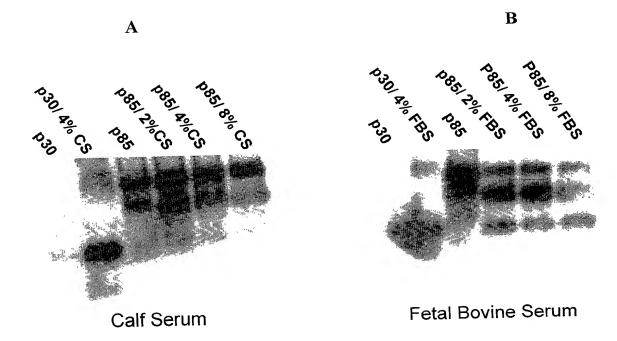
B. 16 KDa

GRSYHDRKSKVDLDRL (SEQ ID NO:24)
GRSYHDRKSKVD...... (SEQ ID NO:25)

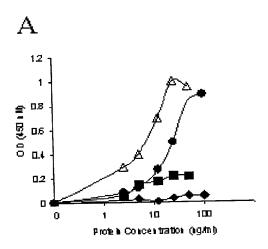
C. 6 Kda

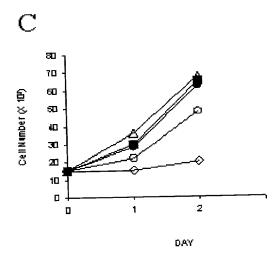


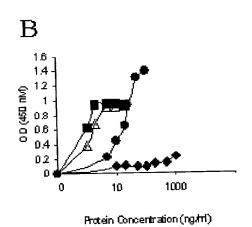
RGRAKTMALVDIQLDHHE (SEQ ID NO:26)
RGRAKTMALVDIQ...... (SEQ ID NO:27)

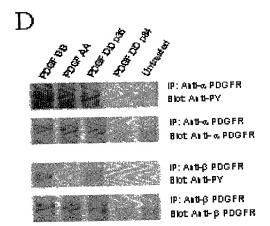


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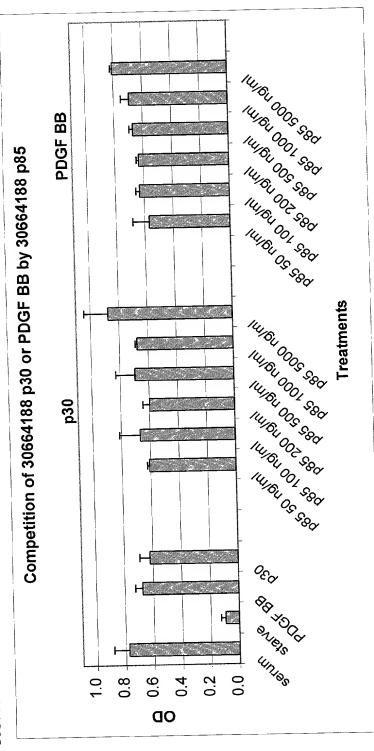


FIG. 18

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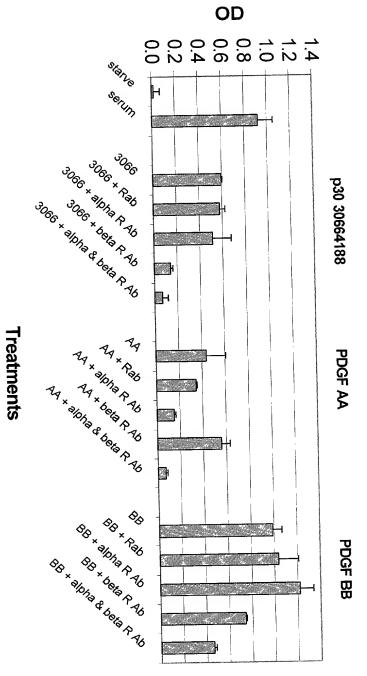
ene fragme	PD GF A	PDGFB	PDGF D
35 17			
22			
154			
z	靣	靣	
8	同		
бľ	同		
69			
0			
z			
τ			
τ			
305			

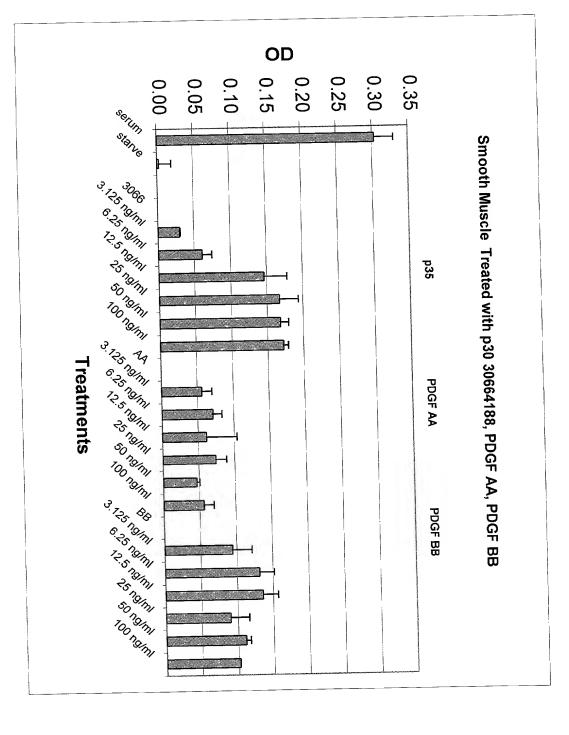
Opt FFIR

FIG. 19

CCD1070 Growth: Competition by Anti-Receptor Antiboldies

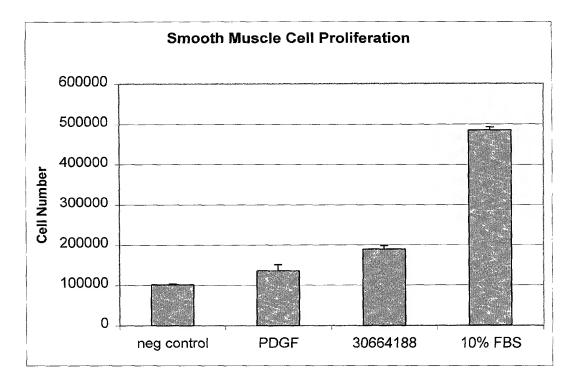
upa e e c





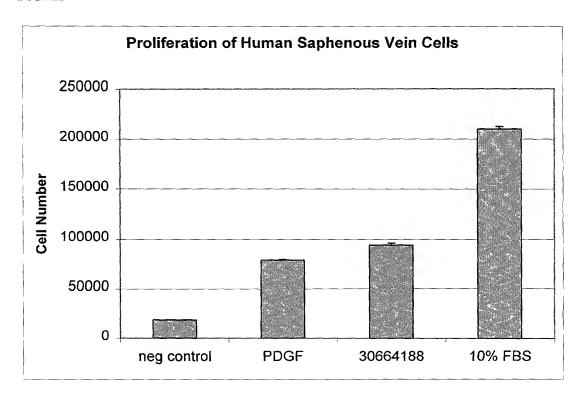
Up t pp t

FIG. 22



tina ma

FIG. 23



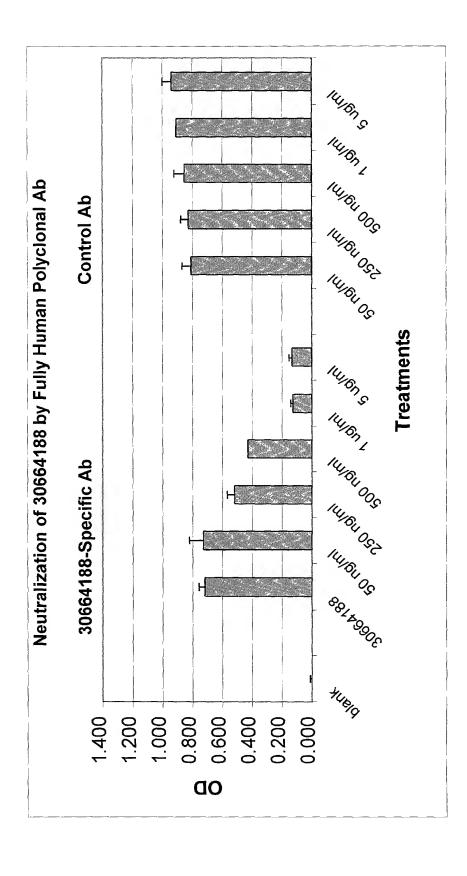


Fig. 25.
Panel A

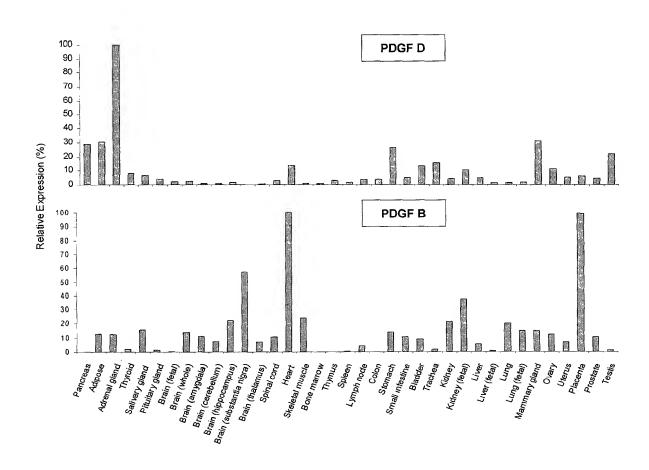
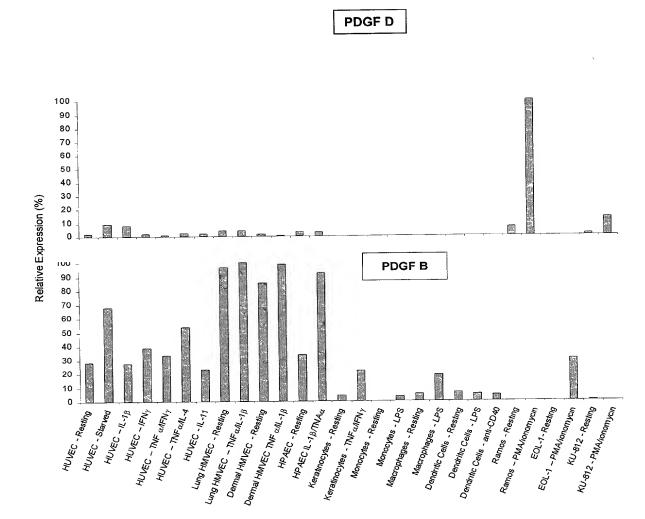


Fig. 25 (cont.)

Panel B



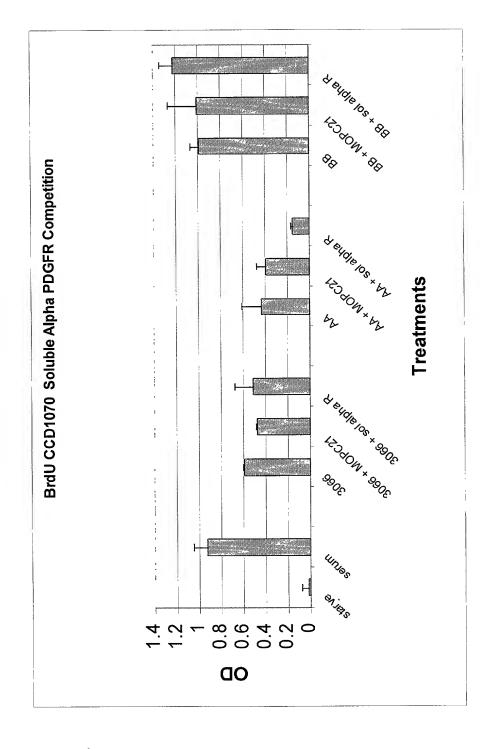
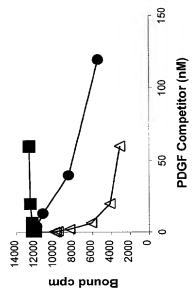
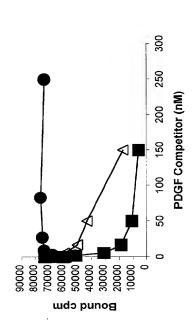


FIG. 27A

FIG. 27B





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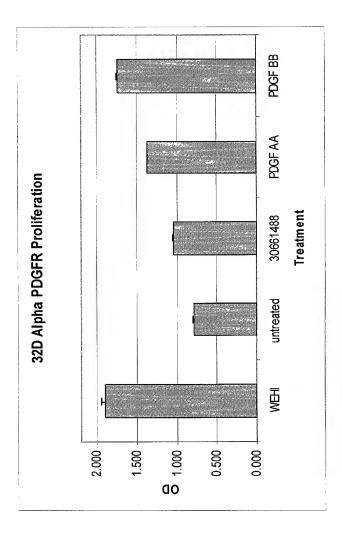


FIG. 29A

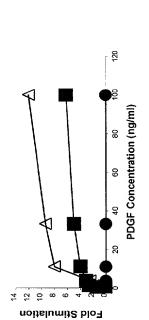


FIG. 29B

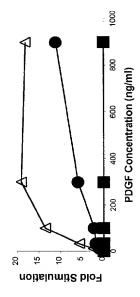


FIG. 29D

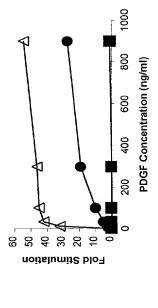
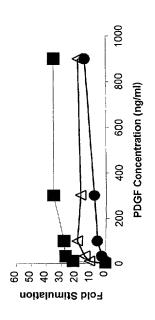


FIG. 29C



IND COMPANY